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Sequence Listing was accepted.

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Reviewer: markspencer

Timestamp: Thu Jul 26 14:09:43 EDT 2007

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Application No: 10591741 Version No: 1.0

Input Set:

Output Set:

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Finished: 2007-07-23 16:37:13.428  
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 420 ms  
Total Warnings: 4  
Total Errors: 0  
No. of SeqIDs Defined: 8  
Actual SeqID Count: 8

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# SEQUENCE LISTING

<110> Meakin, Susan  
Volkening, Kathryn Elizabeth

<120> Method of Proliferating Precursor Cells

<130> 50217/005001

<140> 10591741

<141> 2007-07-23

<150> US 10/591,741

<151> 2006-09-01

<150> PCT/CA05/000345

<151> 2005-03-04

<150> US 60/549,870

<151> 2004-03-04

<160> 8

<170> PatentIn version 3.3

<210> 1

<211> 492

<212> PRT

<213> Homo sapiens

<220>

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<223> FRS3 from human

<400> 1

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Leu	Gly	Ser	Gly	Val	Met	Glu	Leu	Thr	Gln	Ser	Glu	Leu	Val	Leu	His
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Leu	His	Arg	Arg	Glu	Ala	Val	Arg	Trp	Pro	Tyr	Leu	Cys	Leu	Arg	Arg
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Tyr	Gly	Tyr	Asp	Ser	Asn	Leu	Phe	Ser	Phe	Glu	Ser	Gly	Arg	Arg	Cys
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Gln	Thr	Gly	Gln	Gly	Ile	Phe	Ala	Phe	Lys	Cys	Ser	Arg	Ala	Glu	Glu	85	90	95
Ile	Phe	Asn	Leu	Leu	Gln	Asp	Leu	Met	Gln	Cys	Asn	Ser	Ile	Asn	Val	100	105	110
Met	Glu	Glu	Pro	Val	Ile	Ile	Thr	Arg	Asn	Ser	His	Pro	Ala	Glu	Leu	115	120	125
Asp	Leu	Pro	Arg	Ala	Pro	Gln	Pro	Pro	Asn	Ala	Leu	Gly	Tyr	Thr	Val	130	135	140
Ser	Ser	Phe	Ser	Asn	Gly	Cys	Pro	Gly	Glu	Gly	Pro	Arg	Phe	Ser	Ala	145	150	155
Pro	Arg	Arg	Leu	Ser	Thr	Ser	Ser	Leu	Arg	His	Pro	Ser	Leu	Gly	Glu	165	170	175
Glu	Ser	Thr	His	Ala	Leu	Ile	Ala	Pro	Asp	Glu	Gln	Ser	His	Thr	Tyr	180	185	190
Val	Asn	Thr	Pro	Ala	Ser	Glu	Asp	Asp	His	Arg	Arg	Gly	Arg	His	Cys	195	200	205
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Lys	Phe	Val	Leu	Gly	Pro	Thr	Pro	Ala	Arg	Arg	His	Met	Val	Lys	Cys	245	250	255
Gln	Gly	Leu	Cys	Pro	Ser	Leu	His	Asp	Pro	Pro	His	His	Asn	Asn	Asn	260	265	270
Asn	Glu	Ala	Pro	Ser	Glu	Cys	Pro	Ala	Gln	Pro	Lys	Cys	Thr	Tyr	Glu	275	280	285
Asn	Val	Thr	Gly	Gly	Leu	Trp	Arg	Gly	Ala	Gly	Trp	Arg	Leu	Ser	Pro	290	295	300
Glu	Glu	Pro	Gly	Trp	Asn	Gly	Leu	Ala	His	Arg	Arg	Ala	Ala	Leu	Leu			

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Gln	Gln	Leu	Gly	Gly	Glu	Ala	Gly	Asp	Asp	Gly	Asp	Ser	Arg	Asp	Gly
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Ser	Phe	Pro	Val	Pro	Leu	Thr	Arg	Arg	Arg	Gly	Ser	Pro	Arg	Val	Phe
385					390					395					400
Asn	Phe	Asp	Phe	Arg	Arg	Pro	Gly	Pro	Glu	Pro	Pro	Arg	Gln	Leu	Asn
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Tyr	Ile	Gln	Val	Glu	Leu	Lys	Gly	Trp	Gly	Gly	Asp	Arg	Pro	Lys	Gly
			420					425						430	
Pro	Gln	Asn	Pro	Ser	Ser	Pro	Gln	Ala	Pro	Met	Pro	Thr	Thr	His	Pro
			435					440					445		
Ala	Arg	Ser	Ser	Asp	Ser	Tyr	Ala	Val	Ile	Asp	Leu	Lys	Lys	Thr	Val
			450				455				460				
Ala	Met	Ser	Asn	Leu	Gln	Arg	Ala	Leu	Pro	Arg	Asp	Asp	Gly	Thr	Ala
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His Pro Thr Lys Phe Lys Val Thr Asn Val Asp Asp Glu Gly Val Glu  
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Leu Gly Ser Gly Val Met Glu Leu Thr Gln Ser Glu Leu Val Leu His  
35 40 45

Leu His Gln Arg Glu Ala Val Arg Trp Pro Tyr Leu Cys Leu Arg Arg  
50 55 60

Tyr Gly Tyr Asp Ser Asn Leu Phe Ser Phe Glu Ser Gly Arg Arg Cys  
65 70 75 80

Gln Thr Gly Gln Gly Ile Phe Ala Phe Lys Cys Ser Arg Ala Glu Asp  
85 90 95

Ile Phe Asn Leu Leu Gln Asp Leu Met Gln Cys Asn Ser Ile Asn Val  
100 105 110

Thr Glu Glu Pro Val Ile Ile Thr Arg Ser Ser His Pro Pro Glu Leu  
115 120 125

Asp Leu Pro Arg Gly Pro Pro Gln Pro Ala Gly Tyr Thr Val Ser Gly  
130 135 140

Phe Ser Asn Gly Phe Pro Gly Cys Pro Gly Glu Gly Pro Arg Phe Ser  
145 150 155 160

Ala Pro Arg Arg Pro Ser Thr Ser Ser Leu Arg His Pro Ser Pro Gly  
165 170 175

Glu Glu Ser Thr His Thr Leu Ile Ala Ser Glu Glu Gln Ser His Thr  
180 185 190

Tyr Val Asn Thr Pro Thr Gly Asp Glu Asp Gly Arg Ser Arg His Cys  
195 200 205

Leu Gln Pro Leu Pro Glu Gly Arg Val Pro Leu Pro Ala Gln Thr Gln  
210 215 220

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Gln	Ser	Leu	Cys	Pro	Gly	Met	Gln	Asp	Pro	Pro	His	His	Asn	Asn	Asn	260	265	270	
Glu	Gly	Pro	Ser	Glu	Cys	Pro	Ala	Gln	Pro	Lys	Cys	Thr	Tyr	Glu	Asn	275	280	285	
Val	Ser	Gly	Gly	Leu	Gln	Gln	Gly	Ala	Gly	Trp	Arg	Leu	Ser	Pro	Glu	290	295	300	
Glu	Arg	Gly	Trp	Ser	Gly	Leu	Ala	His	Arg	Arg	Ala	Ala	Leu	Leu	His	305	310	315	320
Tyr	Glu	Asn	Leu	Pro	Pro	Leu	Pro	Pro	Val	Trp	Glu	Ser	Gln	Gly	Gln	325	330	335	
Gln	Pro	Gly	Gly	Glu	Ala	Gly	Asp	Asp	Gly	Asp	Ser	Arg	Asp	Gly	Leu	340	345	350	
Thr	Pro	Ser	Ser	Asn	Gly	Phe	Pro	Asp	Gly	Glu	Glu	Asp	Glu	Thr	Pro	355	360	365	
Leu	Gln	Lys	Pro	Thr	Ser	Thr	Arg	Ala	Ser	Ala	Arg	Ser	His	Ser	Gly	370	375	380	
Phe	Pro	Val	Pro	Leu	Thr	Arg	Arg	Arg	Gly	Ser	Pro	Arg	Val	Phe	Asn	385	390	395	400
Phe	Asp	Phe	Arg	Arg	Gln	Gly	Pro	Glu	Pro	Pro	Arg	Gln	Leu	Asn	Tyr	405	410	415	
Ile	Gln	Val	Glu	Leu	Lys	Gly	Trp	Gly	Thr	Ala	Arg	Pro	Lys	Gly	Pro	420	425	430	
Gln	Asn	Pro	Ser	Val	Ser	Gly	Ala	Pro	Gly	Pro	Thr	Pro	His	Pro	Val	435	440	445	

Arg Ser Ser Asp Ser Tyr Ala Val Ile Asp Leu Lys Lys Thr Ala Ala  
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Met Ser Asp Leu Gln Arg Ala Leu Pro Arg Asp Asp Gly Ala Val Arg  
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Lys Thr Arg His Asn Ser Thr Asp Leu Pro Leu  
485 490

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<212> PRT  
<213> Homo sapiens

<220>  
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<223> FRS2 from human

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Leu Gly Ser Gly Ile Met Glu Leu Thr Asp Thr Glu Leu Ile Leu Tyr  
35 40 45

Thr Arg Lys Arg Asp Ser Val Lys Trp His Tyr Leu Cys Leu Arg Arg  
50 55 60

Tyr Gly Tyr Asp Ser Asn Leu Phe Ser Phe Glu Ser Gly Arg Arg Cys  
65 70 75 80

Gln Thr Gly Gln Gly Ile Phe Ala Phe Lys Cys Ala Arg Ala Glu Glu  
85 90 95

Leu Phe Asn Met Leu Gln Glu Ile Met Gln Asn Asn Ser Ile Asn Val  
100 105 110

Val Glu Glu Pro Val Val Glu Arg Asn Asn His Gln Thr Glu Leu Glu  
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Val Pro Arg Thr Pro Arg Thr Pro Thr Thr Pro Gly Phe Ala Ala Gln



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Pro	Val	Gln	Lys	Gln	Leu	Met	Glu	Lys	Glu	Lys	Leu	Glu	Gln	Leu	Gly
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Gly	Tyr	Asp	Ser	Asp	Glu	Arg	Arg	Asp	Ala	Pro	Ser	Val	Asn	Lys	Leu
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Val	Tyr	Glu	Asn	Ile	Asn	Gly	Leu	Ser	Ile	Pro	Ser	Ala	Ser	Gly	Val
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Arg	Arg	Gly	Arg	Leu	Thr	Ser	Thr	Ser	Thr	Ser	Asp	Thr	Gln	Asn	Ile
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Asn	Asn	Ser	Ala	Gln	Arg	Arg	Thr	Ala	Leu	Leu	Asn	Tyr	Glu	Asn	Leu
			340					345					350		
Pro	Ser	Leu	Pro	Pro	Val	Trp	Glu	Ala	Arg	Lys	Leu	Ser	Arg	Asp	Glu
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Asp Asp Asn Leu Gly Pro Lys Thr Pro Ser Leu Asn Gly Tyr His Asn  
370 375 380

Asn Leu Asp Pro Met His Asn Tyr Val Asn Thr Glu Asn Val Thr Val  
385 390 395 400

Pro Ala Ser Ala His Lys Ile Glu Tyr Ser Arg Arg Arg Asp Cys Thr  
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Pro Thr Val Phe Asn Phe Asp Ile Arg Arg Pro Ser Leu Glu His Arg  
420 425 430

Gln Leu Asn Tyr Ile Gln Val Asp Leu Glu Gly Gly Ser Asp Ser Asp  
435 440 445

Asn Pro Gln Thr Pro Lys Thr Pro Thr Thr Pro Leu Pro Gln Thr Pro  
450 455 460

Thr Arg Arg Thr Glu Leu Tyr Ala Val Ile Asp Ile Glu Arg Thr Ala  
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Ala Met Ser Asn Leu Gln Lys Ala Leu Pro Arg Asp Asp Gly Thr Ser  
485 490 495

Arg Lys Thr Arg His Asn Ser Thr Asp Leu Pro Met  
500 505

<210> 4  
<211> 508  
<212> PRT  
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<220>  
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Met Gly Ser Cys Cys Ser Cys Pro Asp Lys Asp Thr Val Pro Asp Asn  
1 5 10 15

His Arg Asn Lys Phe Lys Val Ile Asn Val Asp Asp Asp Gly Asn Glu  
20 25 30

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Thr	Arg	Lys	Arg	Asp	Ser	Val	Lys	Trp	His	Tyr	Leu	Cys	Leu	Arg	Arg	
	50					55					60					
Tyr	Gly	Tyr	Asp	Ser	Asn	Leu	Phe	Ser	Phe	Glu	Ser	Gly	Arg	Arg	Cys	
65					70					75					80	
Gln	Thr	Gly	Gln	Gly	Ile	Phe	Ala	Phe	Lys	Cys	Ala	Arg	Ala	Glu	Glu	
				85					90					95		
Leu	Phe	Asn	Met	Leu	Gln	Glu	Ile	Met	Gln	Asn	Asn	Ser	Ile	Asn	Val	
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Val	Glu	Glu	Pro	Val	Val	Glu	Arg	Ser	Ser	His	Gln	Thr	Glu	Leu	Glu	
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Ser	His	Pro	Ser	Ser	Arg	His	Pro	Ser	Val	Gly	Ser	Ala	Arg	Leu	Pro	
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Ser	Val	Gly	Glu	Glu	Ser	Thr	His	Pro	Leu	Leu	Val	Ala	Glu	Glu	Gln	
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Val	His	Thr	Tyr	Val	Asn	Thr	Thr	Gly	Val	Gln	Glu	Glu	Arg	Lys	Asn	
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Arg	Ala	Ser	Val	His	Val	Pro	Pro	Glu	Ala	Arg	Val	Ser	Asn	Ala	Glu	
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Ser	Asn	Thr	Pro	Lys	Glu	Glu	Pro	Ser	Asn	Pro	Glu	Asp	Arg	Asp	Pro	
225					230				235						240	
Gln	Val	Leu	Leu	Lys	Pro	Glu	Gly	Val	Arg	Phe	Val	Leu	Gly	Pro	Thr	
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Pro	Val	Gln	Lys	Gln	Leu	Met	Glu	Lys	Glu	Lys	Leu	Glu	Gln	Leu	Gly			
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Lys	Asp	Pro	Val	Ser	Gly	Ser	Gly	Ala	Gly	Asn	Thr	Glu	Trp	Asp	Thr			
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Gly	Tyr	Asp	Ser	Asp	Glu	Arg	Arg	Asp	Val	Pro	Pro	Val	Asn	Lys	Leu			
	290					295					300							
Val	Tyr	Glu	Asn	Ile	Asn	Gly	Leu	Ser	Ile	Pro	Ser	Ala	Ser	Gly	Val			
305					310					315					320			
Arg	Arg	Gly	Arg	Leu	Thr	Ser	Thr	Ser	Thr	Ser	Asp	Thr	Gln	Asn	Ile			
				325					330					335				
Asn	Asn	Ser	Ala	Gln	Arg	Arg	Pro	Ala	Leu	Leu	Asn	Tyr	Glu	Asn	Leu			
			340					345					350					
Pro	Ser	Leu	Pro	Pro	Val	Trp	Glu	Ala	Arg	Lys	Leu	Ser	Arg	Asp	Glu			
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Asp	Asp	Asn	Leu	Gly	Pro	Lys	Thr	Pro	Ser	Leu	Asn	Gly	Tyr	His	Asn			
	370					375					380							
Asn	Leu	Asp	Pro	Met	His	Asn	Tyr	Val	Asn	Thr	Glu	Asn	Val	Thr	Val			
385					390					395					400			
Pro	Ala	Ser	Ala	His	Lys	Ile	Asp	Tyr	Ser	Lys	Arg	Arg	Asp	Cys	Thr			
				405					410					415				
Pro	Thr	Val	Phe	Asn	Phe	Asp	Ile	Arg	Arg	Pro	Ser	Leu	Glu	His	Arg			
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Gln	Leu	Asn	Tyr	Ile	Gln	Val	Asp	Leu	Glu	Gly	Gly	Ser	Asp	Ser	Asp			
	435						440					445						
Asn	Pro	Gln	Thr	Pro	Lys	Thr	Pro	Thr	Thr	Pro	Leu	Pro	Gln	Thr	Pro			
	450					455					460							
Thr	Arg	Arg	Thr	Glu	Leu	Tyr	Ala	Val	Ile	Asp	Ile	Glu	Arg	Thr	Ala			
465					470					475					480			
Ala	Met	Ser	Asn	Leu	Gln	Lys	Ala	Leu	Pro	Arg	Asp	Asp	Gly	Thr	Ser			

485

490

495

Arg Lys Thr